HematoVision - Project Report

Title

HematoVision: Advanced Blood Cell Classification Using Transfer Learning

Objective

To develop a deep learning-based web application capable of classifying blood cell images into their respective types (Eosinophils, Lymphocytes, Monocytes, Neutrophils) using transfer learning techniques, improving diagnostic speed and reliability in hematological analysis.

Technologies Used

- Language: Python

- Frameworks: TensorFlow, Keras, Flask

- Frontend: HTML, CSS

- Backend: Flask (app.py)

- Libraries: NumPy, Matplotlib, PIL, OS

- Model: MobileNetV2 (pre-trained CNN)

- Dataset: Blood Cell Images Dataset (12,000+ images)

Project Directory (C:/hem2)

├── static/

SmartBridge (AI-ML Internship)

SmartBridge (AI-ML Internship)

- Real-time prediction shown with image preview.

How to Run

1. Train the Model (first time only):

python train_model.py

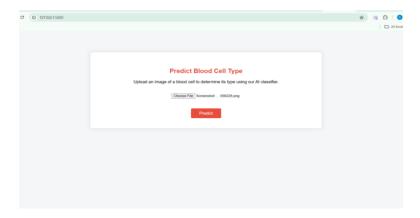
2. Launch the Web App:

python app.py

Then open: http://127.0.0.1:5000 in your browser.

Output

- Upload any blood cell image.



- The app displays the predicted cell type along with the uploaded image.



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Applications

- Assisting pathologists in automated blood cell classification.
- Can be extended to detect abnormalities or other cell types in the future.

Learning Outcomes

- Real-world application of Transfer Learning in Medical AI.
- Full-stack integration using Flask.
- Experience with image classification pipelines and deployment.